

10 20 30 40
TPVSEKQLAE VVANTITPLM KAQSVPGMAV AVIYQGKPHY

50 60 70 80 90 100
YTFGKADIAA NKPVTPQTLF ELGSISKTFT GVLGGDAIAR GEISLDDAVT RYWPQLTGKQ

110 120 130 140 150 160
WQGIRMLDLA TYTAGGLPLQ VPDEVTDNAS LLRFYQNWQP QWKPGTTRLY ANASILGFGA

170 180 190 200 210 220
LAVKPSGMPY EQAMTTRVLK PLKLDHTWIN VPKAEEAHYA WGYRDGKAVR VSPGMLDAQA

230 240 250 260 270 280
YGVKTNVQDM ANWVMANMAP ENVADASLKQ GIALAQSRYW RIGSMYQGLG WEMLNWPVEA

290 300 310 320 330 340
NTVVEGSDSK VALAPLPVAE VNPPAPPVKA SWVHKTGSTG GFGSYVAFIP EKOQIGIVMLA

350 360
NTSYPNPARV EAAYHILEAL Q

FIG. 1

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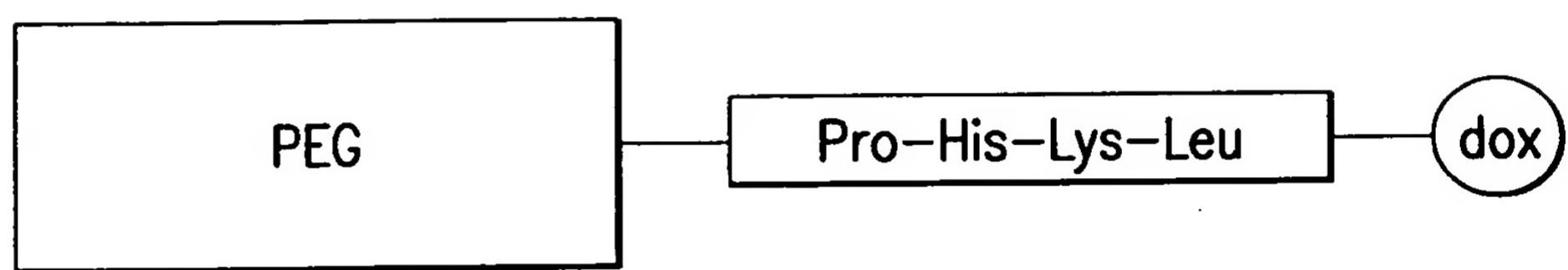


FIG. 2

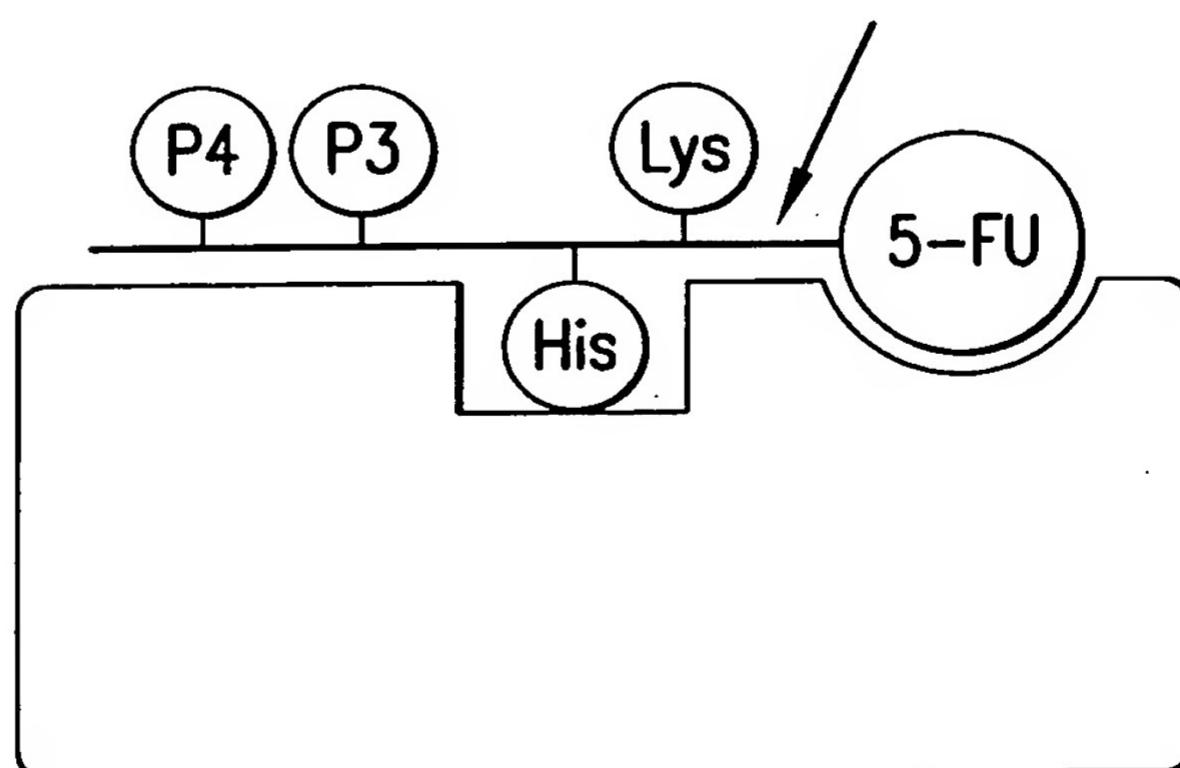


FIG. 3

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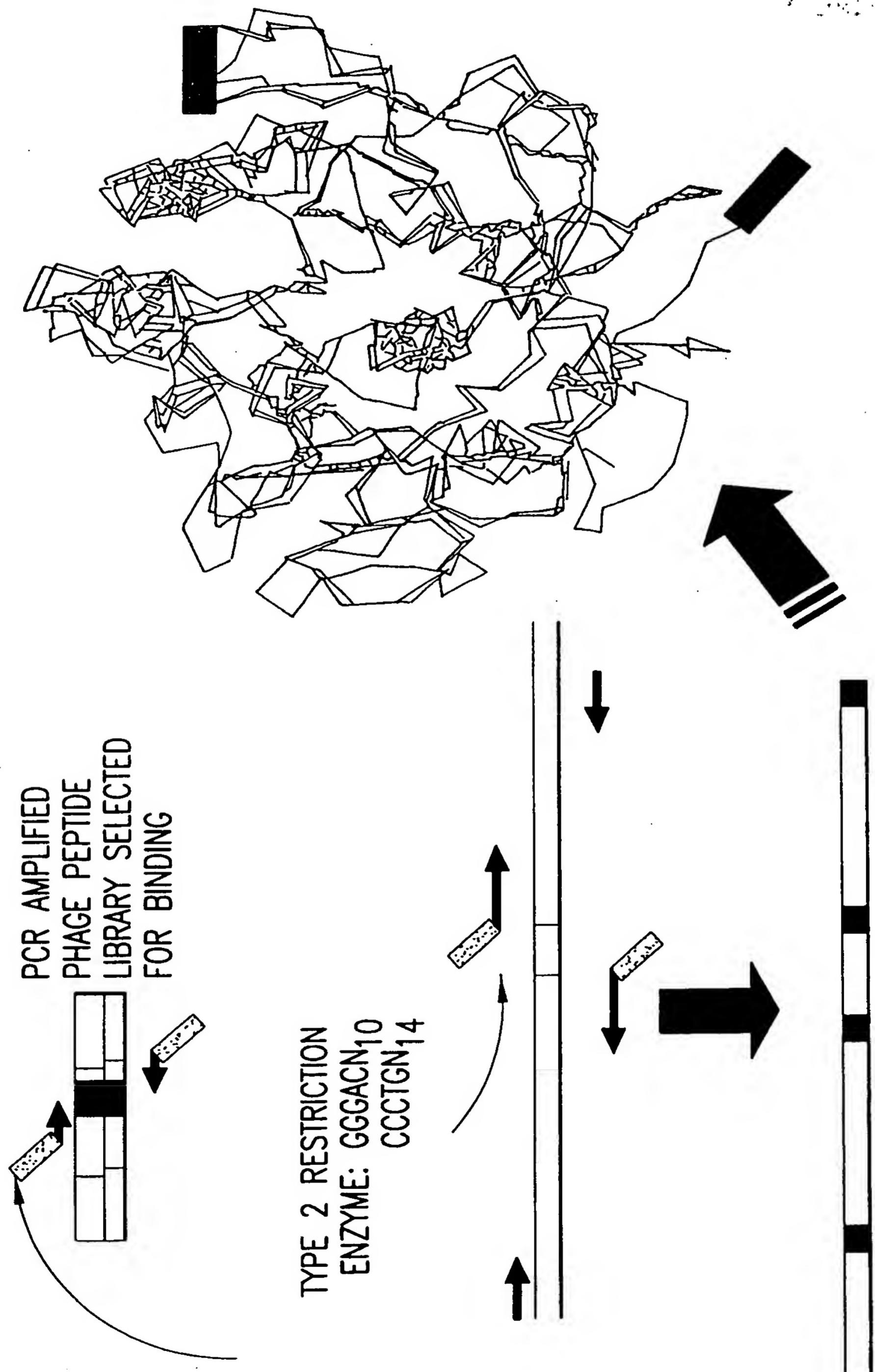
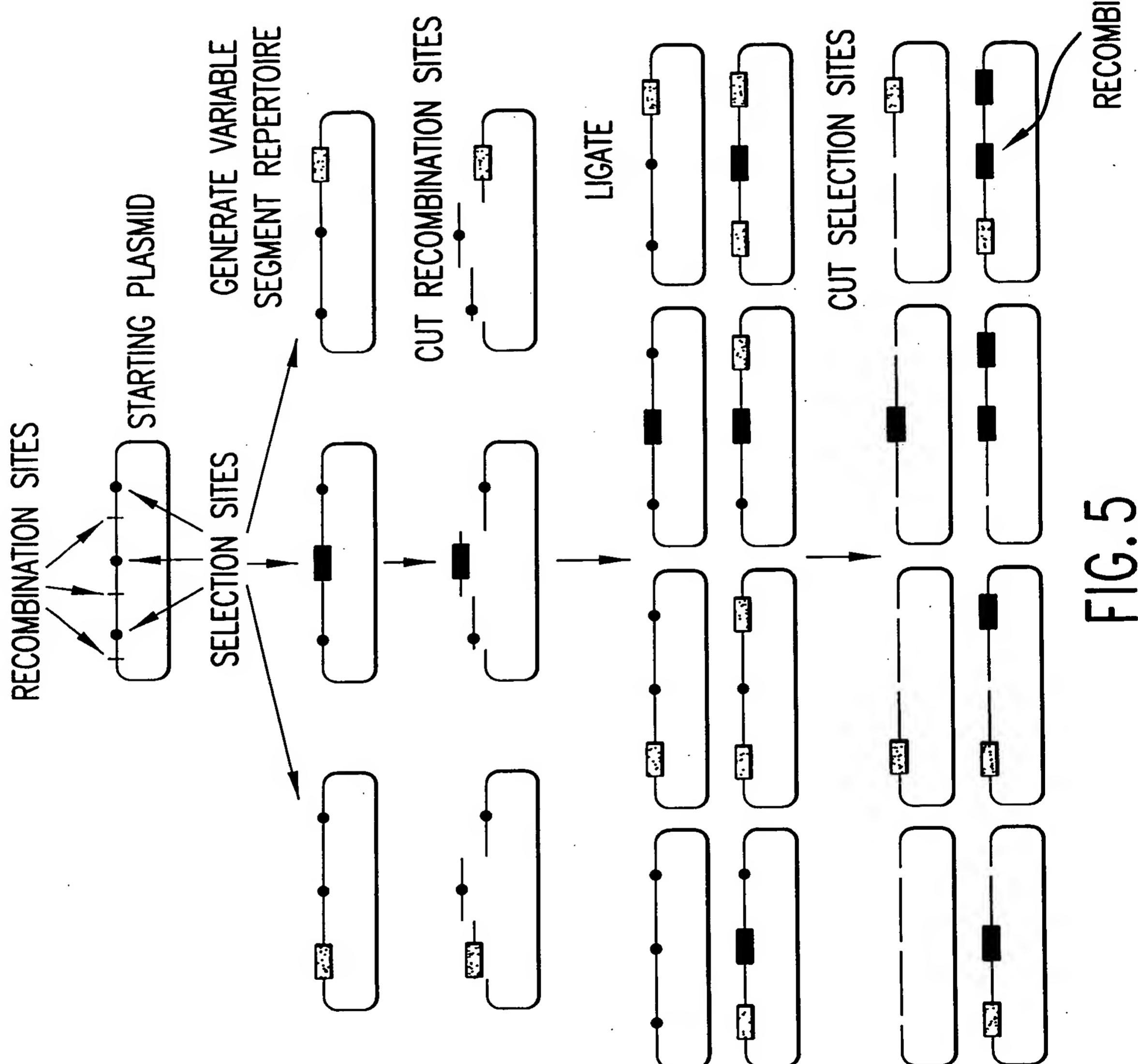


FIG. 4



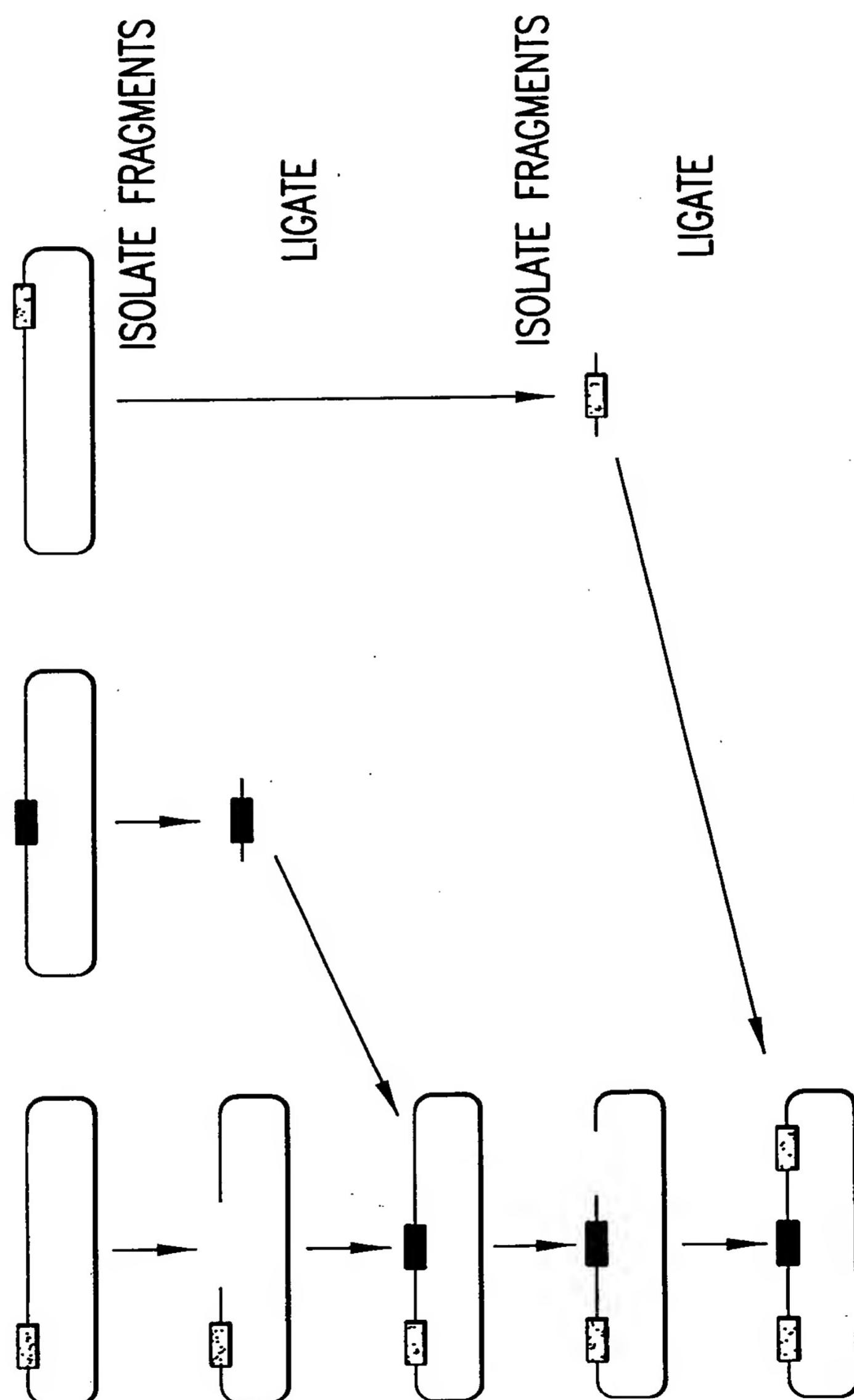


FIG. 6

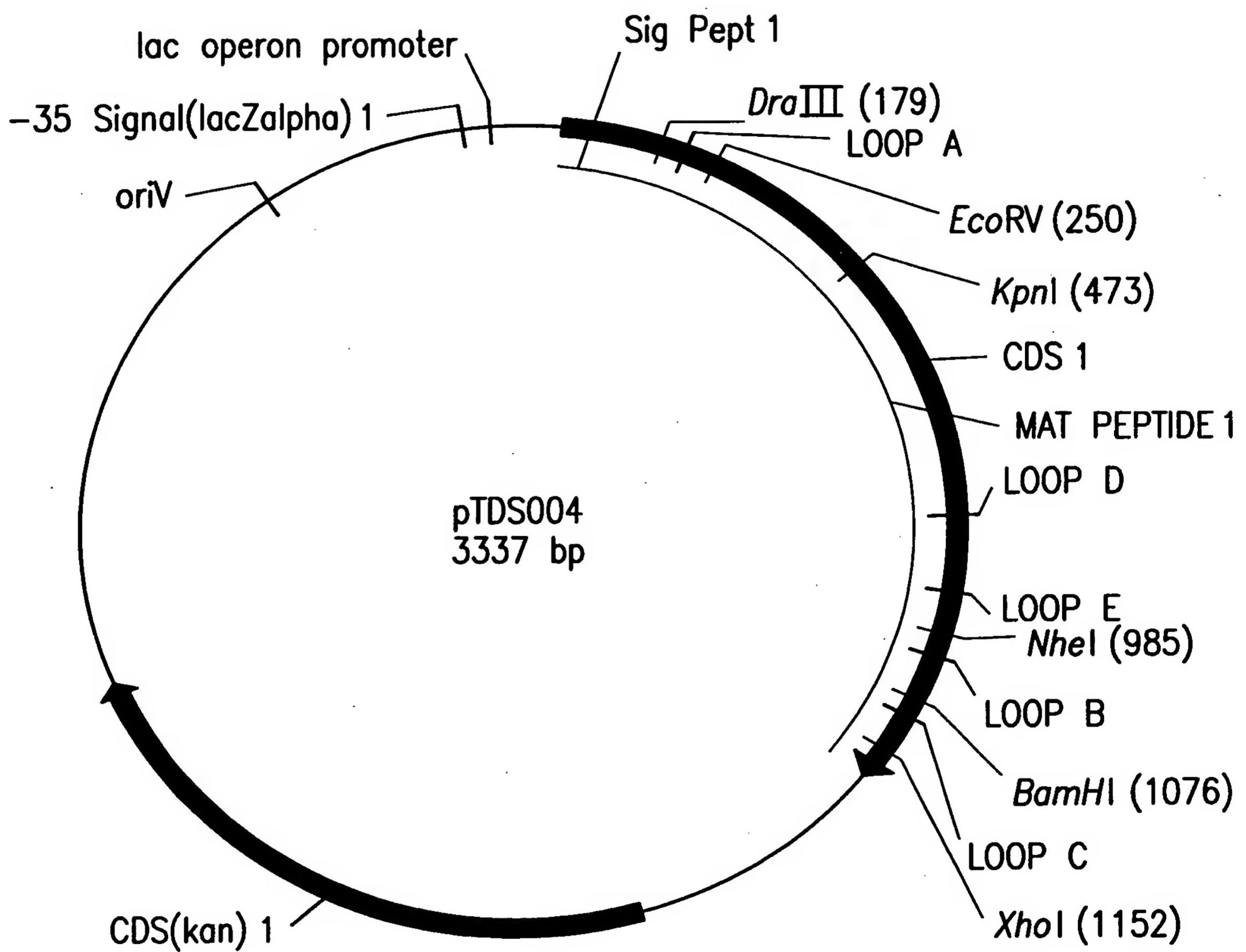


FIG. 7

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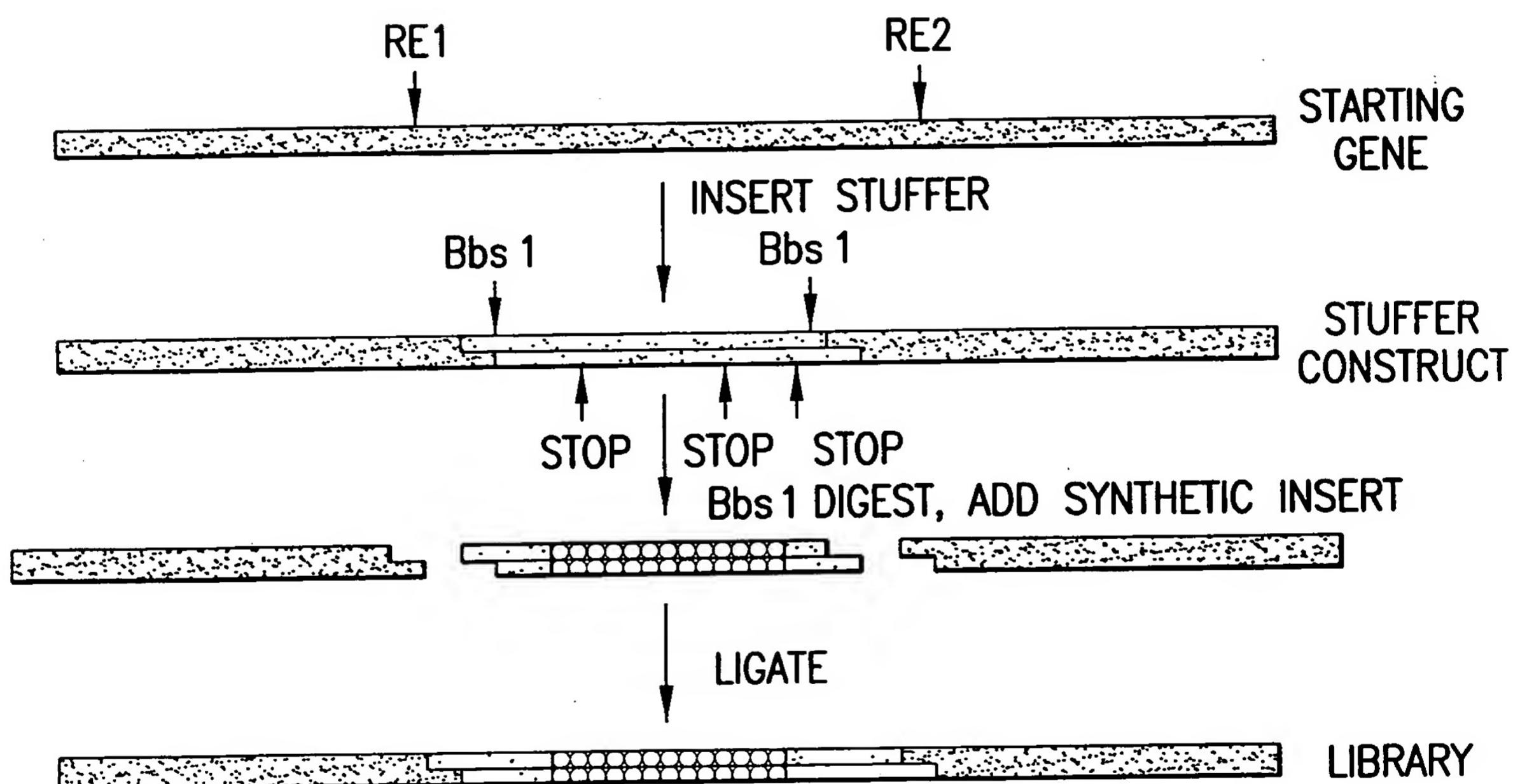


FIG. 8

